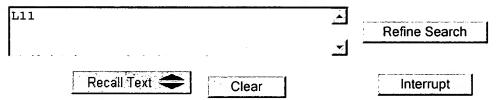
Refine Search

Search Results -

Terms	Documents
L7 same treat\$3 same L5	6

US Pre-Grant Publication Full-Text Database
US Patents Full-Text Database
US OCR Full-Text Database
EPO Abstracts Database
JPO Abstracts Database
Derwent World Patents Index
IBM Technical Disclosure Bulletins

Search:



Search History

DATE: Monday, November 07, 2005 Printable Copy Create Case

Set Name side by side	Query	<u>Hit</u> <u>Count</u>	Set Name result set
DB=B	PGPB, USPT, USOC, EPAB, JPAB, DWPI; PLUR = YES; OP = OR		
<u>L11</u>	L7 same treat\$3 same L5	6	<u>L11</u>
<u>L10</u>	L7 with treat\$3 with L5	2	<u>L10</u>
<u>L9</u>	6190412	. 10	<u>L9</u>
<u>L8</u>	L7 and L5	63	<u>L8</u>
<u>L7</u>	(cartilage adj growth) or (bone adj growth) or (skeletal adj growth)	8470	<u>L7</u>
<u>L6</u>	L1 and L5	15	<u>L6</u>
<u>L5</u>	L4 or L3 or L2	11091	<u>L5</u>
<u>L4</u>	ODF or (osteoclast adj differentiation adj factor)	1154	<u>L4</u>
<u>L3</u>	RANKL or (receptor adj activator adj of adj NF-kB adj ligand)	626	<u>L3</u>
<u>L2</u>	TRANCE or (tumor adj necrosis adj factor adj related adj activation adj induced adj cytokine)	9548	<u>L2</u>
<u>L1</u>	acromegaly or gigantism or (exostosis adj bursata) or (exostosis adj cartilaginea) or (multiple adj osteocartilaginous adj exostoses)	2088	<u>L1</u>

FILE 'MEDLINE, BIOSIS, EMBASE, SCISEARCH, CAPLUS, USPATFULL, PCTFULL' ENTERED AT 17:47:53 ON 07 NOV 2005 L1 3816 S TRANCE OR (TUMOR(W) NECROSIS(W) FACTOR(W) RELATED(W) ACTIVATION(W L26804 S RANKL OR (RECEPTOR(W)ACTIVATOR(W)OF(W)NFKB(W)LIGAND) L3 4906 S ODF OR (OSTEOCLAST(W) DIFFERENTIAT? (W) FACTOR) 12655 S L1 OR L2 OR L3 L4L5129051 S CARTILAGE(S)(GROWTH OR DIFFERENTIATION) OR BONE(S)GROWTH 28106 S ACROMEGALY OR GIGANTISM OR (EXOSTOSIS(W)BURSATA) OR (EXOSTOSI L6 L7 1818 S (THERAP? OR TREAT? OR ADMINISTER?) (S) L4 L8192 S L7(P)(L5 OR L6) L9 80 S L7(S)(L5 OR L6) L10 71 DUP REM L9 (9 DUPLICATES REMOVED) L11161 DUP REM L8 (31 DUPLICATES REMOVED) L12 6 S L10 AND 1960-2001/PY L13 16 S L11 AND 1960-2001/PY L14 226 S L4(S)(L5 OR L6) L15 · 167 DUP REM L14 (59 DUPLICATES REMOVED) L16 23 S L15 AND 1960-2001/PY

GenCore version 5.1.6 Copyright (c) 1993 - 2005 Compugen Ltd.

OM protein - protein search, using sw model

Run on: November 3, 2005, 17:40:04; Search time 44 Seconds

(without alignments)

325.741 Million cell updates/sec

Title: US-09-933-915A-3

Perfect score: 1014

Sequence: 1 AVQKELQHIVGSQHIRAEKA.....LLDPDQDATYFGAFKVRDID 192

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 513545 seqs, 74649064 residues

Total number of hits satisfying chosen parameters: 513545

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database : Issued_Patents_AA:*

1: /cgn2_6/ptodata/1/iaa/5A_COMB.pep:*

2: /cgn2_6/ptodata/1/iaa/5B_COMB.pep:*

3: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*

4: /cgn2_6/ptodata/1/iaa/6B_COMB.pep:*
5: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*

6: /cgn2_6/ptodata/1/iaa/backfiles1.pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

		ક				
Result		Query				
No.	Score	Match	Length	DB	ID ·	Description
1	1014	100.0	317	3	US-08-996-139-13	Sequence 13, Appl
2	1014	100.0	317	3	US-08-995-659-13	Sequence 13, Appl
3	1014	100.0	317	3	US-09-215-649A-13	Sequence 13, Appl
4	1014	100.0	317	3	US-09-052-521C-4	Sequence 4, Appli
5	1014	100.0	317	4	US-09-577-780-13	Sequence 13, Appl
6	1014	100.0	317	4	US-09-577-800-13	Sequence 13, Appl
7	1014	100.0	317	4	US-09-466-496-13	Sequence 13, Appl
8	1014	100.0	317	4	US-09-871-856-13	Sequence 13, Appl
9	1014	100.0	317	4	US-09-871-291-13	Sequence 13, Appl
10	1014	100.0	317	4	US-09-396-937-2	Sequence 2, Appli
11	1014	100.0	317	4	US-09-877-650-13	Sequence 13, Appl
12	1014	100.0	317	4	US-09-865-363-13	Sequence 13, Appl
13	1014	100.0	317	4	US-09-688-459-13	Sequence 13, Appl
14	877	86.5	294	3	US-08-996-139-11	Sequence 11, Appl
15	877	86.5	294	3	US-08-995-659-11	Sequence 11, Appl
16	877	86.5	294	3	US-09-215-649A-11	Sequence 11, Appl
17	877	86.5	294	4	US-09-577-780-11	Sequence 11, Appl
						

```
877
18
            86.5
                   294 4 US-09-577-800-11
                                                     Sequence 11, Appl
      877
                   294 4 US-09-466-496-11
19
            86.5
                                                     Sequence 11, Appl
20
      877
            86.5
                   294 4 US-09-871-856-11
                                                     Sequence 11, Appl
21
      877
            86.5
                   294 4 US-09-871-291-11
                                                     Sequence 11, Appl
22
      877
            86.5
                   294 4 US-09-877-650-11
                                                     Sequence 11, Appl
                                                     Sequence 11, Appl
23
      877
            86.5
                   294 4 US-09-865-363-11
      877
                                                     Sequence 11, Appl
24
            86.5
                   294 4 US-09-688-459-11
      877
            86.5
25
                   316 2 US-08-842-842-7
                                                     Sequence 7, Appli
      877
            86.5
26
                   316 3 US-08-989-362-2
                                                     Sequence 2, Appli
            86.5
      877
27
                   316 3 US-09-052-521C-2
                                                     Sequence 2, Appli
28
      877
            86.5
                   316 4 US-09-671-658A-2
                                                     Sequence 2, Appli
29
      877
            86.5
                   316 4 US-09-396-937-4
                                                     Sequence 4, Appli
                                                     Sequence 6, Appli
      877
            86.5
                   316 4 US-09-396-937-6
30
                   187 4 US-09-396-937-8
31
      771
            76.0
                                                     Sequence 8, Appli
                   173 4 US-09-396-937-10
            75.8
32
      769
                                                     Sequence 10, Appl
                 173 4 US-09-396-937-12
33
      759
            74.9
                                                     Sequence 12, Appl
            71.2 188 4 US-09-396-937-14
    721.5
34
                                                     Sequence 14, Appl
    711.5
            70.2 182 4 US-09-396-937-16
35
                                                     Sequence 16, Appl
            68.1 173 4 US-09-396-937-18
36
      691
                                                     Sequence 18, Appl
            66.1 173 4 US-09-396-937-20
37
      670
                                                     Sequence 20, Appl
      418
          41.2 77 4 US-09-632-287A-11
38
                                                     Sequence 11, Appl
39
      363
          35.8
                   77 4 US-09-632-287A-10
                                                     Sequence 10, Appl
40
    232.5
          22.9 161 4 US-09-565-423-7
                                                     Sequence 7, Appli
41
    232.5
            22.9 253 3 US-09-320-424-11
                                                     Sequence 11, Appl
            22.9
42
    232.5
                   253 4 US-09-825-563-11
                                                     Sequence 11, Appl
            22.9
22.9
43
    232.5
                   256 3 US.-09-320-424-13
                                                     Sequence 13, Appl
                   256 4 US-09-825-563-13
279 3 US-09-072-993C-3
    232.5
44
           22.9
22.9
                                                     Sequence 13, Appl
45
    232.5
                                                     Sequence 3, Appli
```

ALIGNMENTS

```
RESULT 2
US-08-995-659-13
; Sequence 13, Application US/08995659
; Patent No. 6242213
; GENERAL INFORMATION:
    APPLICANT: Anderson, Dirk M.
    APPLICANT: Galibert, Laurent
    APPLICANT: Maraskovsky, Eugene
    TITLE OF INVENTION: Ligand for Receptor Activator of NF-kappaB NUMBER OF SEQUENCES: 19
    CORRESPONDENCE ADDRESS:
      ADDRESSEE: Immunex Corporation, Law Department
      STREET: 51 University Street
      CITY: Seattle
      STATE: WA
      COUNTRY: USA
      ZIP: 98101
    COMPUTER READABLE FORM:
      MEDIUM TYPE: Floppy disk
      COMPUTER: Apple Power Macintosh
      OPERATING SYSTEM: Apple Operating System 7.5.5
      SOFTWARE: Microsoft Word for Power Macintosh 6.0.1
    CURRENT APPLICATION DATA:
      APPLICATION NUMBER: US/08/995,659
      FILING DATE: 22 DECEMBER 1997
       CLASSIFICATION:
     PRIOR APPLICATION DATA:
      APPLICATION NUMBER: USSN 60/064,671
      FILING DATE: 14 OCTOBER 1997
      CLASSIFICATION:
```

```
PRIOR APPLICATION DATA:
     APPLICATION NUMBER: USSN 08/813,509
     FILING DATE: 07 MARCH 1997
     CLASSIFICATION:
    PRIOR APPLICATION DATA:
     APPLICATION NUMBER: USSN 08/772,330
     FILING DATE: 23 DECEMBER 1996
     CLASSIFICATION:
   ATTORNEY/AGENT INFORMATION:
     NAME: Perkins, Patricia Anne
     REGISTRATION NUMBER: 34,693
     REFERENCE/DOCKET NUMBER: 2852-A
    TELECOMMUNICATION INFORMATION:
     TELEPHONE: (206)587-0430
     TELEFAX: (206)233-0644
  INFORMATION FOR SEQ ID NO: 13:
    SEQUENCE CHARACTERISTICS:
     LENGTH: 317 amino acids
     TYPE: amino acid
     TOPOLOGY: linear
    MOLECULE TYPE: protein
US-08-995-659-13
 Query Match
                     100.0%; Score 1014; DB 3; Length 317;
 Best Local Similarity 100.0%; Pred. No. 3.2e-103;
 Matches 192; Conservative 0; Mismatches 0; Indels
                                                     0; Gaps
                                                                 0;
          1 AVOKELOHIVGSOHIRAEKAMVDGSWLDLAKRSKLEAOPFAHLTINATDIPSGSHKVSLS 60
Qy
            126 AVQKELQHIVGSQHIRAEKAMVDGSWLDLAKRSKLEAQPFAHLTINATDIPSGSHKVSLS 185
Db
         61 SWYHDRGWAKISNMTFSNGKLIVNQDGFYYLYANICFRHHETSGDLATEYLQLMVYVTKT 120
Qy
            186 SWYHDRGWAKISNMTFSNGKLIVNODGFYYLYANICFRHHETSGDLATEYLOLMVYVTKT 245
Db
Qy
        121 SIKIPSSHTLMKGGSTKYWSGNSEFHFYSINVGGFFKLRSGEEISIEVSNPSLLDPDQDA 180
           246 SIKIPSSHTLMKGGSTKYWSGNSEFHFYSINVGGFFKLRSGEEISIEVSNPSLLDPDQDA 305
Db
Qу
        181 TYFGAFKVRDID 192
            Db
        306 TYFGAFKVRDID 317
```